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Entry Information

Entry from: EMBL

Entry Options

Launch analysis tool:
BlastN

Link to related information:

Save entry:

View:

ID AA565544 standard; mRNA; EST; 457 BP.
 XX
 AC AA565544;
 XX
 SV AA565544.1
 XX
 DT 11-SEP-1997 (Rel. 52, Created)
 DT 03-MAR-2000 (Rel. 62, Last updated, Version 2)
 XX
 DE nk42d05.s1 NCI_CGAP_GC2 Homo sapiens cDNA clone IMAGE:1016
 DE sequence.
 XX
 KW EST.
 XX
 OS Homo sapiens (human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.
 XX
 RN [1]
 RP 1-457
 RA NCI-CGAP;
 RT "National Cancer Institute, Cancer Genome Anatomy Project
 RT Gene Index http://www.ncbi.nlm.nih.gov/ncicgap";
 RL Unpublished.
 XX
 DR RZPD; IMAGp998J102536.
 DR RZPD; IMAGp998J10253606.
 DR UNILIB; 939; 939.
 XX
 CC Insert Length: 1022 Std Error: 0.00
 CC Seq primer: -40m13 fwd. ET from Amersham
 CC High quality sequence stop: 426.
 XX
 CC On Sep 12, 1996 this sequence version replaced gi:1393368.
 CC Contact: Robert Strausberg, Ph.D.
 CC Tel: (301) 496-1550
 CC Email: Robert_Strausberg@nih.gov
 CC Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael
 CC Emmert-Buck, M.D., Ph.D.
 CC cDNA Library Preparation: Stratagene, Inc., David B. Krizm
 CC Ph.D.
 CC cDNA Library Arraying: Greg Lennon, Ph.D.
 CC DNA Sequencing by: Washington University Genome Sequencing
 CC Clone distribution: NCI-CGAP clone distribution information
 CC found through the I.M.A.G.E. Consortium/LLNL at:
 CC www-bio.llnl.gov/bbrp/image/image.html
 XX
 FH Key Location/Qualifiers
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 FT source 1..457
 FT /db_xref="taxon:9606"
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 FT /db_xref="RZPD: IMAGp998J102536"
 FT /db_xref="RZPD: IMAGp998J102536"
 FT /db_xref="RZPD: IMAGp998J10253606"

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Launch analysis tool:
BlastN

Link to related information:

Save entry:

View:

ID AA813484 standard; mRNA; EST; 422 BP.
 XX
 AC AA813484;
 XX
 SV AA813484.1
 XX
 DT 16-FEB-1998 (Rel. 54, Created)
 DT 03-MAR-2000 (Rel. 62, Last updated, Version 3)
 XX
 DE ai67b05.s1 Soares_testis_NHT Homo sapiens cDNA clone 13758
 DE sequence.
 XX
 KW EST.
 XX
 OS Homo sapiens (human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.
 XX
 RN [1]
 RP 1-422
 RA NCI-CGAP;
 RT "National Cancer Institute, Cancer Genome Anatomy Project
 Gene Index <http://www.ncbi.nlm.nih.gov/ncicgap>";
 RL Unpublished.
 XX
 DR RZPD; [IMAGp998E103473](#).
 DR UNILIB; 628; 628.
 XX
 CC Insert Length: 1355 Std Error: 0.00
 CC Seq primer: -40m13 fwd. ET from Amersham
 CC High quality sequence stop: 403.
 XX
 CC On Sep 29, 1997 this sequence version replaced gi:1520753.
 CC Contact: Robert Strausberg, Ph.D.
 CC Tel: (301) 496-1550
 CC Email: Robert_Strausberg@nih.gov
 CC cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatim
 CC Bonaldo, Ph.D.
 CC cDNA Library Arrayed by: Greg Lennon, Ph.D.
 CC DNA Sequencing by: Washington University Genome Sequencing
 CC Clone distribution: NCI-CGAP clone distribution informatio
 CC found through the I.M.A.G.E. Consortium/LLNL at:
 CC www-bio.llnl.gov/bbrp/image/image.html
 XX
 FH Key Location/Qualifiers
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 FT source 1..422
 FT /db_xref="taxon:9606"
 FT /db_xref="RZPD: [IMAGp998E103473](#)"
 FT /db_xref="RZPD: [IMAGp998E103473](#)"
 FT /db_xref="RZPD: [IMAGp998E103473](#)"
 FT /db_xref="UNILIB: 628"
 FT /mol_type="mRNA"
 FT /note="Vector: pT7T3D-Pac (Pharmacia) with
 FT polylinker; Site_1: Not I; Site_2: Eco RI;
 FT was prepared from mRNA obtained from Clont

FT /db_xref="RZPD: IMAGp998J10253606"
FT /db_xref="RZPD: IMAGp998J10253606"
FT /db_xref="RZPD: IMAGp998J10253606"
FT /db_xref="UNILIB: 939"
FT /mol_type="mRNA"
FT /note="Vector: Bluescript SK-; Site_1: Eco
FT Cloned unidirectionally. Primer: Oligo dT
FT tumor. 5' adaptor sequence: 5' GAATTCCGGCAC
FT sequence: 5' CTCGAGTTTTTTTTTTTTTTTT 3' A
FT size: 1.2 kb."
FT /organism="Homo sapiens"
FT /clone="IMAGE:1016169"
FT /clone_lib="NCI_CGAP_GC2"
FT /tissue_type="germ cell tumor"
FT /lab_host="SOLR (kanamycin resistant)"
XX
SQ Sequence 457 BP; 104 A; 99 C; 135 G; 119 T; 0 other;
tttgaaaatc aaacatttaa taaaactgtgg ggctgggggt ggggagaact tgc
ggggagcacg aagcctgggc ctatggagag ggctggactt gctacatttt ggg
cacaaggagc tttgggaaaa ttgttttttgc tgccaagatg cattatagaa cca
caatttctca gtgtggccca aatggtggcc tcacagtctt ctcctggaga ctg
ggcgtgagtc tcccgccagc tgtcagcatt caccatctt gttgtctgt act
cagtggcttg agaagtagtt tcagccacag aagaaacgaa cacgtctggg ggc
tgccgggacg tggtggggac ttcccttag agtggtctgg ccccatctg agt
tctgctcagg ctggagacac aggtcttgtt acacaaa
//

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Entry Information

Entry from: EMBL

Entry Options

Launch analysis tool:

BlastN

Link

Link to related information:

Save entry:

View:

ID HS1749 standard; mRNA; EST; 175 BP.

XX

AC T19174;

XX

SV T19174.1

XX

DT 20-DEC-1994 (Rel. 42, Created)

DT 04-MAR-2000 (Rel. 63, Last updated, Version 4)

XX

DE h01003s Testis 2 Homo sapiens cDNA clone h01003 5' end, mR

XX

KW EST.

XX

OS Homo sapiens (human)

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele

OC Eutheria; Primates; Catarrhini; Hominidae; Homo.

XX

RN [1]

RP 1-175

RX MEDLINE; 95301283.

RX PUBMED; 7782076.

RA Pawlak A., Toussaint C., Levy, I., Bulle F., Poyard M., Bar

RA Guellaen G.;

RT "Characterization of a large population of mRNAs from huma

RL Genomics 26(1):151-158(1995).

XX

DR UNILIB; 120; 120.

XX

CC Contact: Guellaen G

CC Unite INSERM 99

CC INSERM

CC Unite INSERM 99, Hopital Henri Mondor, 94010 Creteil, Fran

CC Tel: (33)149813530

CC Fax: (33)148980908

CC Email: guellaen@infobiogen.fr

CC Seq primer: SP6.

XX

FH Key Location/Qualifiers

FH

FT source 1..175

FT /db_xref="taxon:9606"

FT /db_xref="UNILIB:120"

FT /mol_type="mRNA"

FT /note="Vector: PGEM 5zf(+); Site_1: Sal1; mRNA was prepared from human testis of a 2 cDNA was prepared using a 15mer oligo dT a degenerated bases at its 3'end and contain at its 5'end. The cDNA was cloned between sites of PGEM 5zf(+). The 3' end is at the Sal1 site is lost during the cloning proce corresponding to abundant species were eli library."

FT /organism="Homo sapiens"

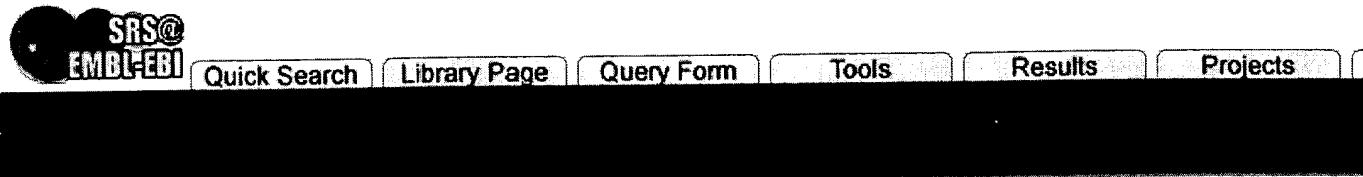
FT /clone="h01003"

FT /clone_lib="Testis 2"

XX

SO Sequence 175 bp. 16 A. 27 C. 59 G. 22 T. 2 other.

52 sequence 1/5 D5; 40 A, 37 C, 30 G, 22 T; 4 others,
tacggaaatgg agcgacaggg agtggacgtg ccccatgtga aatgcaaaga cca
cagcccttgg gggagagcaa ggagcatccg cggatgggaa gagaactgcg agg
tggtggaggg ccagctagtg ccanttcca gctgacggtc ctggaagnga agt
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Reset

Entry Information	
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Entry Options	
Launch analysis tool:	
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Link to related information:	
<input type="button" value="Link"/>	
Save entry: <input type="button" value="Save"/>	
View: <input type="button" value="Printer Friendly"/>	

aaatcccttcc tggccccc tttttttt tttttttt tttttttt tttttttt
tctgcctcca aattttctgt tcagattctg agaaccttca tctgagggct gtn
ttggggcata agtggcatgg tccttctggc t
//


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Entry from: <u>EMBL</u>	
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Save entry:	
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View:	
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ID MM1276610 standard; mRNA; EST; 394 BP.
 XX
 AC AA386755;
 XX
 SV AA386755.1
 XX
 DT 25-JUN-1997 (Rel. 52, Created)
 DT 03-MAR-2000 (Rel. 62, Last updated, Version 2)
 XX
 DE vc17h03.r1 Ko mouse embryo 11 5dpc Mus musculus cDNA clone
 DE mRNA sequence.
 XX
 KW EST.
 XX
 OS Mus musculus (house mouse)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
 OC Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
 XX
 RN [1]
 RP 1-394
 RA Marra M., Hillier L., Allen M., Bowles M., Dietrich N., Du
 RA Geisel S., Kucaba T., Lacy M., Le M., Martin J., Morris M.
 RA Schellenberg K., Steptoe M., Tan F., Underwood K., Moore B
 RA Wylie T., Lennon G., Soares B., Wilson R., Waterston R.;
 RT "The WashU-HHMI Mouse EST Project";
 RL Unpublished.
 XX
 DR RZPD; [IMAGp998D061908](#).
 DR UNILIB; [845](#); 845.
 XX
 CC On Sep 12, 1996 this sequence version replaced gi:1288550.
 CC Contact: Marra M/Mouse EST Project
 CC WashU-HHMI Mouse EST Project
 CC Washington University School of MedicineP
 CC 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
 CC Tel: 314 286 1800
 CC Fax: 314 286 1810
 CC Email: mouseest@watson.wustl.edu
 CC This clone is available royalty-free through LLNL ; contac
 CC IMAGE Consortium (info@image.llnl.gov) for further informa
 CC MGI:467725
 CC High quality sequence stop: 388.
 XX
 FH Key Location/Qualifiers
 FH
 FT source 1..394
 FT /db_xref="taxon:10090"
 FT /db_xref="RZPD: [IMAGp998D061908](#)"
 FT /db_xref="RZPD: [IMAGp998D061908](#)"
 FT /db_xref="RZPD: [IMAGp998D061908](#)"
 FT /db_xref="RZPD: [IMAGp998D061908](#)"
 FT /db_xref="UNILIB: [845](#)"
 FT /mol_type="mRNA"
 FT /note="Organ: embryo; Vector: pSPORT1; Site_1: NotI; Site_2: NotI; Total RNAs were extracted from embryos (excluding placenta and yolk sac)"
 FT

FT double-stranded cDNA was synthesized with
FT primer GAGAGAGACTAGTTCTAGATCGCGAGCGGCCGCTT
FT 3'. The cDNAs were ligated to LL-Sal3A:
FT GCTATTGACGTCGACTATCC 3' and LL-Sal3B: 5'
FT GGATAGTCGACGTCAAT 3'. The cDNAs were size
FT amplified by long-range PCR using Ex Taq p
FT cycles. The PCR-amplifiable cDNA mixture
FT round of equalization and was digested wi
FT cloned into the SalI/NotI sites of the pS
FT vector (Life Technologies). The library w
FT by Dr. Minoru S. H. Ko and Dr. Xiaohong Wa
FT /sex="pooled"
FT /organism="Mus musculus"
FT /strain="C57BL/6J"
FT /clone_lib="Ko mouse embryo 11 5dpc"
FT /dev_stage="11.5dpc"
FT /lab_host="DH10B"
FT /clone="IMAGE:774869"
FT /tissue_type="embryo"
XX
SQ Sequence 394 BP; 124 A; 97 C; 110 G; 61 T; 2 other;
ctatccaaaa gacggaatgg agcgccacgg agtggcctcc cgtggagtt aag
agccgcgan nttgtggaga gcggagagca tcggcagagt gagaaccacg agg
cggctcagtgc ccccgagtg ccagttgcca gctgccagga cccttctcct ctc
tagcattgaa accctgaaga agaaagccca ggaactgatt gaaaacatca atg
gcagaaggac catgcactta tgaccaactt cagggacagc ctcaagatga agg
tctgacagaa aagttggagg agaggatgtt ccaggtgtac agccaccaca gca
tcagggaaaga ctacaagaat ttacccagaa gatg
//

FT Inc., and primed with a Not I - oligo(dT)
FT TGTTACCAATCTGAAGTGGGAGCGGGCCGCCAATTTTTTT
FT Double-stranded cDNA was ligated to Eco RI
FT (Pharmacia), digested with Not I and clone
FT and Eco RI sites of the modified pT7T3 vec
FT through one round of normalization to Cot5
FT constructed by Bento Soares and M. Fatima
FT /sex="male"
FT /organism="Homo sapiens"
FT /clone="1375857"
FT /clone_lib="Soares_testis_NHT"
FT /lab_host="DH10B"
XX
SQ Sequence 422 BP; 93 A; 90 C; 126 G; 113 T; 0 other;
ttttgaaaa tcaaacatt aataaactgt ggggctgggg gtggggagaa ctt
gtggggagca cgaaggcctgg gcctatggag cgggctggac ttgctacatt ttg
agcacaagga gctttgggtt tttgttttt tctgccaaga tgcattatag aac
aacaatttct cagtgtggcc caaatggtagg cctcacagtc ttctcctggaa gac
aaggcgtgag tctcccgga gctgtcagca ttcaccatct ctgttggct gta
agcagttggcc tgagaagtag tttcagccac agaagaaacg aacacgtctg ggg
gttgcgggaa cgtgggtgggg actttcccta gagtggctcg gccccatctg agt
tc
//